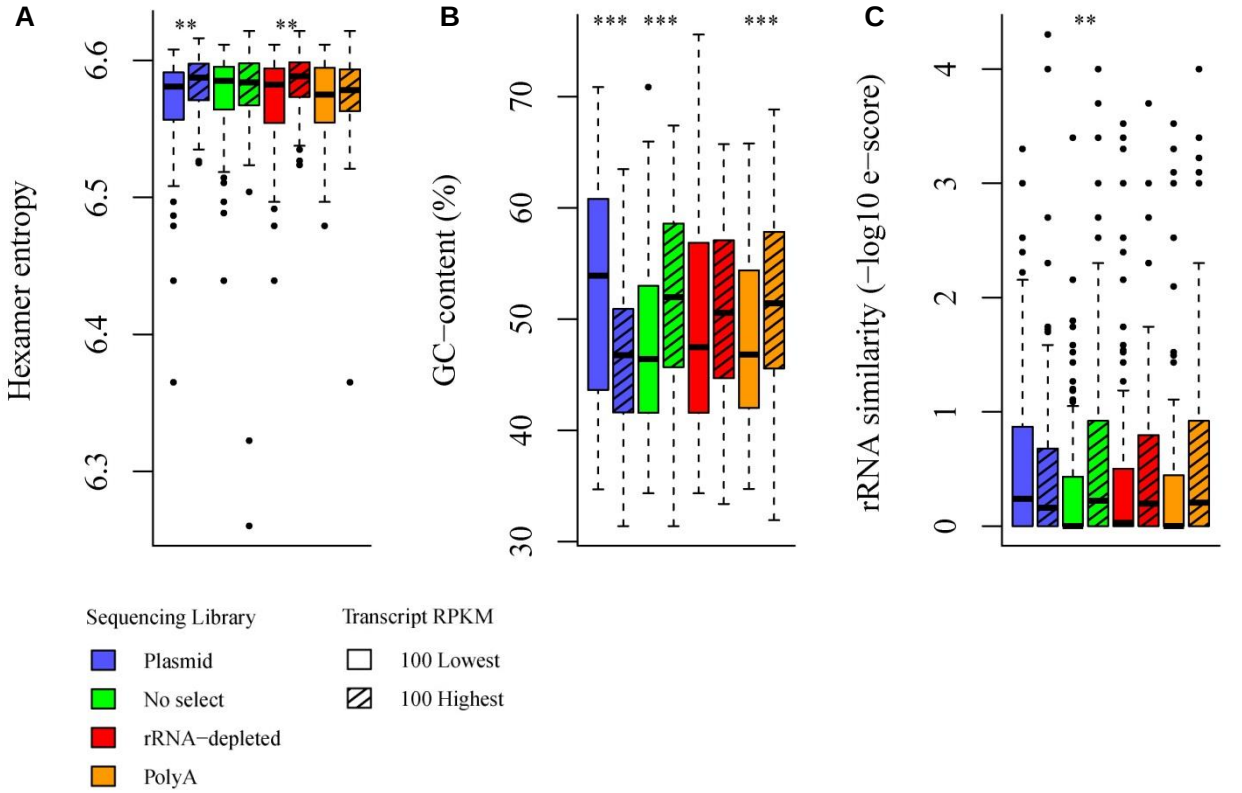
Figure S8


A
Hexamer entropy
6.3
6.4
6.5
6.6
**
**
B
GC−content (%)
30
40
50
60
70
***
***
***
C
rRNA similarity (−log10 e−score)
0
1
2
3
4
**
Sequencing Library
Plasmid
No select
rRNA−depleted
PolyA
Transcript RPKM
100 Lowest
100 Highest